> **This Raw Listing contains the General Information Section and up to first 5 pages.**

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 1   <110> APPLICANT: Probst, Peter
 2         Bhatia, Ajay
 3         Skeiky, Yasir
 4         Fling, Steve
 5         Maisonneuve, Jeff
 6   <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND
 7         DIAGNOSIS OF CHLAMYDIAL INFECTION
 8   <130> FILE REFERENCE: 210121.469C4
 9   <140> CURRENT APPLICATION NUMBER: US/09/454,684
10   <141> CURRENT FILING DATE: 1999-12-03
11   <160> NUMBER OF SEQ ID NOS: 303
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20         caaaataaga actctgcttt catgcagcct gtgaacgtat ccgctgattt agctgccatc    180
21         gttggtgcag gacctatgcc tcgcacagag atcattaaga aaatgtggga ttacattaag    240
22         gagaatagtc ttcaagatcc tacaaacaaa cgtaatatca atcccgatga taaattggct    300
23         aaagtttttg gaactgaaaa acctatcgat atgttccaaa tgacaaaaat ggtttctcaa    360
24         cacatcatta aataaaatag aaattgactc acgtgttcct cgtctttaag atgaggaact    420
25         agttcattct ttttgttcgt ttttgtgggt attactgtat ctttaacaac tatcttagca    480
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28   <211> LENGTH: 183
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34         gctaaagttt ttggaactga aaaacctatc gatatgttcc aaatgacaaa aatggtttct    180
35         caa                                                                  183
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37   <211> LENGTH: 110
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50         tagggccagc tctttctaaa gagattattg ctagattgca gttgaatccc gaagctagag   180
51         ctgcagagtt gactgaggaa gaggttggtc gactaaacgc tcttttacag tcggattacg   240
52         ttgttgaagg ggatttgcgc cgtcgtgtgc aatctgatat caaacgtctg attactatcc   300
53         atgcttatcg tggacaaaga catagacttt ctttgcctgt tcgtggtcag agaacaaaaa   360
54         caaattctcg cacgcgtaag ggtaaacgta aaactattgc aggtaagaag aaataataat   420
55         ttttaggaga gagtgttttg gttaaaaatc aagcgcaaaa aagaggcgta aaaagaaaac   480
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57         taaccataac agacc                                                    555
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65         Ala Asp Leu Ala Ala Ile Val Gly Ala Gly Pro Met Pro Arg Thr Glu
66                     20                  25                  30
67         Ile Ile Lys Lys Met Trp Asp Tyr Ile Lys Glu Asn Ser Leu Gln Asp
68                 35                  40                  45
69         Pro Thr Asn Lys Arg Asn Ile Asn Pro Asp Asp Lys Leu Ala Lys Val
70             50                  55                  60
71         Phe Gly Thr Glu Lys Pro Ile Asp Met Phe Gln Met Thr Lys Met Val
72         65                  70                  75                  80
73         Ser Gln His Ile Ile Lys
74                         85
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76    <211> LENGTH: 61
77    <212> TYPE: PRT
78    <213> ORGANISM: Chlamydia trachomatis
79    <400> SEQUENCE: 6
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83                     20                  25                  30
84         Asn Ile Asn Pro Asp Asp Lys Leu Ala Lys Val Phe Gly Thr Glu Lys
85                 35                  40                  45
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87             50                  55                  60
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89    <211> LENGTH: 36
90    <212> TYPE: PRT
91    <213> ORGANISM: Chlamyida trachomatis
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 97          Gln Ser His Arg
 98                  35
 99     <210> SEQ ID NO 8
100     <211> LENGTH: 18
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108     <211> LENGTH: 5
109     <212> TYPE: PRT
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131                  35
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140                      20                  25                  30
141          Ile Ile Ala Arg Leu Gln Leu Asn Pro Glu Ala Arg Ala Ala Glu Leu
142                  35                  40                  45
143          Thr Glu Glu Glu Val Gly Arg Leu Asn Ala Leu Leu Gln Ser Asp Tyr
144                  50                  55                  60
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145        Val Val Glu Gly Asp Leu Arg Arg Arg Val Gln Ser Asp Ile Lys Arg
146         65                  70                  75                  80
147        Leu Ile Thr Ile His Ala Tyr Arg Gly Gln Arg His Arg Leu Ser Leu
148                             85                  90                  95
149        Pro Val Arg Gly Gln Arg Thr Lys Thr Asn Ser Arg Thr Arg Lys Gly
150                            100                 105                 110
151        Lys Arg Lys Thr Ile Ala Gly Lys Lys Lys
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154    <211> LENGTH: 20
155    <212> TYPE: PRT
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163    <211> LENGTH: 20
164    <212> TYPE: PRT
165    <213> ORGANISM: Chlamydia trachomatis
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172    <211> LENGTH: 161
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186        attaaggttg ccaagtctgc tgccgaattg accgcaaata ttttggaaca agctggaggc    180
187        gcgggctctt ccgcacacat tacagcttcc caagtgtcca aaggattagg ggatgcgaga    240
188        actgttgtcg ctttagggaa tgcctttaac ggagcgttgc caggaacagt tcaaagtgcg    300
189        caaagcttct tctctcacat gaaagctgct agtcagaaaa cgcaagaagg ggatgagggg    360
190        ctcacagcag atctttgtgt gtctcataag cgcagagcgg ctgcggctgt ctgtagcatc    420
191        atcggaggaa ttacctacct cgcgacattc ggagctatcc gtccgattct gtttgtcaac    480
192        aaaatgctgg caaaaccgtt tctttcttcc caaactaaag caaatatggg atcttctgtt    540
193        agctatatta tggcggctaa ccatgcagcg tctgtggtgg gtgctggact cgctatcagt    600
194        gcggaaagag cagattgcga agcccgctgc gctcgtattg cgagagaaga gtcgttactc    660
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207                        20                  25                  30
208        Lys Thr Lys Gly Met Asp Lys Thr Ile Lys Val Ala Lys Ser Ala Ala
209                35                  40                  45
210        Glu Leu Thr Ala Asn Ile Leu Glu Gln Ala Gly Gly Ala Gly Ser Ser
211             50                  55                  60
212        Ala His Ile Thr Ala Ser Gln Val Ser Lys Gly Leu Gly Asp Ala Arg
213        65                  70                  75                  80
214        Thr Val Val Ala Leu Gly Asn Ala Phe Asn Gly Ala Leu Pro Gly Thr
215                        85                  90                  95
216        Val Gln Ser Ala Gln Ser Phe Phe Ser His Met Lys Ala Ala Ser Gln
217                        100                 105                 110
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219                115                 120                 125
220        His Lys Arg Arg Ala Ala Ala Ala Val Cys Ser Ile Ile Gly Gly Ile
221             130                 135                 140
222        Thr Tyr Leu Ala Thr Phe Gly Ala Ile Arg Pro Ile Leu Phe Val Asn
223        145                 150                 155                 160
224        Lys Met Leu Ala Lys Pro Phe Leu Ser Ser Gln Thr Lys Ala Asn Met
225                        165                 170                 175
226        Gly Ser Ser Val Ser Tyr Ile Met Ala Ala Asn His Ala Ala Ser Val
227                        180                 185                 190
228        Val Gly Ala Gly Leu Ala Ile Ser Ala Glu Arg Ala Asp Cys Glu Ala
229                        195                 200                 205
230        Arg Cys Ala Arg Ile Ala Arg Glu Glu Ser Leu Leu Glu Val Pro Gly
231             210                 215                 220
232        Glu Glu Asn Ala Cys Glu Lys Lys Val Ala Gly Glu Lys Ala Lys Thr
233        225                 230                 235                 240
234        Phe Thr Arg Ile Lys Tyr Ala Leu Leu Thr Met Leu Glu Lys Phe Leu
235                        245                 250                 255
236        Glu Cys Val Ala Asp Val Phe Lys Leu Val Pro Leu Pro Ile Thr Met
237                        260                 265                 270
238        Gly Ile Arg Ala Ile Val Ala Ala Gly Cys Thr Phe Thr Ser Ala Ile
239                275                 280                 285
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241             290                 295
242    <210> SEQ ID NO 18
243    <211> LENGTH: 18
       <212> TYPE: PRT
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Input Set: **I454684.RAW**

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620	W	"N" or "Xaa" used: Feature required	gatccgaatt cggcacgagn tttttcctgt tttttctt
645	W	"N" or "Xaa" used: Feature required	ccgacaacgt attcattacg tgtaggcggt ttagaaag
1014	W	"N" or "Xaa" used: Feature required	gatccgaatt cggcacgaga aggtagatcc gatntcag
1092	W	"N" or "Xaa" used: Feature required	gaaaccttgn tctcttcgnc tctctcctgt agcanaca
1093	W	"N" or "Xaa" used: Feature required	tcagcgtatt cggactgatg ccctaaagat cccnggan
1122	W	"N" or "Xaa" used: Feature required	aagtattttt atagctgaag aacataggat gagtcatg
1123	W	"N" or "Xaa" used: Feature required	atgaaagact gaataagcta tttgatagcc cctttagt
1124	W	"N" or "Xaa" used: Feature required	nagctnagaa caaaattgct agaggagatg ttcgttct
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1958	W	"N" or "Xaa" used: Feature required	ctcgtagcag atctttgtgt gtctcataag cgcanagc
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3473	W	"N" or "Xaa" used: Feature required	Gly Ser Lys Val Xaa Phe
4668	W	"N" or "Xaa" used: Feature required	Phe Phe Thr Gly Asn Ser Ala Thr Asn Gly G
4672	W	"N" or "Xaa" used: Feature required	Gln Xaa Thr Leu Phe Ala Ser Asn Ser Ala L
4694	W	"N" or "Xaa" used: Feature required	Leu Lys Asp Arg Ala Val Leu Ser Xaa Pro S
4696	W	"N" or "Xaa" used: Feature required	Gln Ala Leu Leu Ile Met Glu Ala Gly Thr S
4698	W	"N" or "Xaa" used: Feature required	Asp Leu Lys Leu Xaa Thr Xaa Ser Ile Pro L
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6211	W	"N" or "Xaa" used: Feature required	gcgnaaagag cagattgcga agcccgctgc gctcgtat
6249	W	"N" or "Xaa" used: Feature required	Val Gly Ala Gly Leu Ala Ile Ser Ala Xaa A
6324	W	"N" or "Xaa" used: Feature required	tctctgttac agataaggag actagangca tctagttt
6549	W	"N" or "Xaa" used: Feature required	cgtgccgnt

PREVIOUSLY ERRORED SEQUENCES-EDITED

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 1    <210> 141
 2    <211> 18
 3    <212> PRT
 4    <213> Artificial Sequence
 5    <220>
 6    <223> Made in a lab
 7    <400> 141
 8          Tyr Leu Ala Thr Phe Gly Ala Ile Arg Pro Ile Leu Phe Val Asn Lys
 9          1                   5                   10                  15
10          Met Leu
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> This Raw Listing contains the General
> Information Section and those Sequences
> containing ERRORS.

Does Not Comply
Corrected Diskette Needed

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 1   <110> Probst, Peter
 2         Bhatia, Ajay
 3         Skeiky, Yasir
 4         Fling, Steve
 5         Maisonneuve, Jeff
 6   <120> COMPOSITIONS AND METHODS FOR TREATMENT AND
 7         DIAGNOSIS OF CHLAMYDIAL INFECTION
 8   <130> 210121.469C4
 9   <140> US/09/454,684
10   <141> 1999-12-03
11   <160> 303
12   <170> FastSEQ for Windows Version 3.0/4.0
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ERRORED SEQUENCES FOLLOW

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       13   <210> 141
       14   <211> 18
       15   <212> PRT
       16   <213> Artificial Sequence
       17   <220>
       18   <223> Made in a lab
E-->   19   <400> 141
       20         Tyr Leu Ala Thr Phe Gly Ala Ile Arg Pro Ile Leu Phe Val Asn Lys
       21         1               5                   10                  15
       22         Met Leu
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Line	?	Error/Warning	Original Text
19	E	Sequence IDs 14 and 141 differ	<400> 14

OIPE 02Co

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/454,684

CRF Processing Date: 12/20/99
Edited by: _____
Verified by: _____ (STIC staff)

ENTERED

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically:
_____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
_____

☑ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: 141

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included:
_____

☐ Deleted extra, invalid, headings used by an applicant, specifically:
_____

☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____ .

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically:
_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically:
_____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other:
_____
_____
_____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95